## IN THE CLAIMS

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Please cancel Claims 1 through 38, 50, and 51.

Please amend the claims as indicated in Appendix A as attached hereto. A clean version of the claims follow:

- 39. (Amended) A cell that produces a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, the cell comprising heterologous genes encoding one or more sugar-nucleotide regenerating enzyme and one or more glycosyltransferase.
- 49. (Amended) A method of producing a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, comprising the step of contacting a cell comprising heterologous genes encoding:
  - (i). one or more encoding sugar-nucleotide regenerating enzymes selected from the group consisting of GalK, GalT, GalU,
    PykF, Ndk, PpK, AcK, PoxB, Ppa, PgM, NagE, Agm1, glmU, a GalNAc kinase, a pyrophosphorylase, Ugd, NanA, Cmk, NeuA, Alg2, Alg1, SusA, ManB, ManC, a phosphomannomutase,
    GalE, GMP, GMD, and GFS; and
  - (ii). one or more glycosyltransferase,

with a bioenergetic.

Please add the following new claims:

- 52. (New) The cell of claim 39, wherein the one or more sugar-nucleotide regenerating enzyme is selected from the group consisting of GalK, GalT, GalU, PykF, Ndk, PpK, AcK, PoxB, Ppa, PgM, NagE, Agm1, glmU, a GalNAc kinase, a pyrophosphorylase, Ugd, NanA, Cmk, NeuA, Alg2, Alg1, SusA, ManB, ManC, a phosphomannomutase, GalE, GMP, GMD, and GFS.
- 53. (New) The cell of claim 39 comprising genes encoding GalK, GalT, and GalU.
  - 54. (New) The cell of claim 53 comprising a gene encoding Ndk.
  - 55. (New) The cell of claim 53 comprising a gene encoding Ppk.
  - 56. (New) The cell of claim 53 comprising a gene encoding PykF.
- 57. (New) The cell of claim 53 comprising genes encoding PoxB, Ndk, and Ppa.
  - 58. (New) The cell of claim 39 comprising a gene encoding SusA.
  - 59. (New) The cell of claim 58 further comprising a gene encoding GalE.
  - 60. (New) The cell of claim 58 further comprising a gene encoding GluT.
- 61. (New) The cell of claim 58 further comprising genes encoding Ugd and UGT2B7.

- 70. (New) The cell of claim 62, wherein the fucosyltransferase is selected from the group consisting of  $\alpha$ 1,3-FucT,  $\alpha$ 1,2-FucT, and  $\alpha$ 1,3/4-FucT.
- 71. (New) A method of producing a glycoconjugate of interest in the absence of an exogenously supplied nucleotide triphosphate, the method comprising:

contacting a cell comprising heterologous genes encoding susA, galE, and a glycosyltransferase with a bioenergetic.

- 72. (New) The method according to claim 71, wherein the bioenergetic comprises fructose generated within the cell by susA acting on sucrose.
- 73. (New) The method according to claim 72 further comprising supplying the sucrose to the cell.
- 74. (New) The method according to claim 71, wherein the glycosyltransferase is a galactosyltransferase.
- 75. (New) The method according to claim 74, wherein the galactosyltransferase is  $\alpha 1,3 GalT$ .
- 76. (New) The method according to claim 71, wherein the glycosyltransferase is LgtC.

- 62. (New) The cell of claim 39. wherein the one more glycosyltransferase(s) is selected from the group consisting galactosyltransferase, a glucosyltransferase, a N-acetylglucosaminyl transferase, an N-acetylgalactosaminyl transferase, a glucuronyltransferase, a sialyltransferase, a mannosyltransferase, and a fucosyltransferase.
- 63. (New) The cell of claim 62 wherein the galactosyltransferase is selected from the group consisting of LgtB and LgtC.
- 64. (New) The cell of claim 62, wherein the glucosyltransferase is selected from the group consisting of LgtF, Alg5, and DUGT.
- 65. (New) The cell of claim 62, wherein the N-acetylglucosaminyl transferase is LgtA.
- 66. (New) The cell of claim 62, wherein the N-acetylgalactosaminyl transferase is UDP-GalNAc:2'-fucosylgalactoside-α-3-*N*-acetylgalactosaminyl transferase.
- 67. (New) The cell of claim 62, wherein the glucuronyltransferase is UGT2B7.
  - 68. (New) The cell of claim 62, wherein the sialyltransferase is SiaT 0160.
- 69. (New) The cell of claim 62, wherein the mannosyltransferase is selected from the group consisting of Alg1 and Alg2.